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48kD i-antigen pro
48 kDa immobilizat
IAG48 (G1) surface
55kD i-antigen pro
55 kDa immobilizat
Synthetic 55kD i-a
48kD i-antigen re
48kD i-antigen re
48kD i-antigen re
                                                                                                                                              February 11, 2003, 19:43:19; Search time 32.0571 Seconds (without alignments) 1837.243 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1: /SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980_DAT:*
2: /SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981_DAT:*
3: /SIDS2/gcgdata/geneseqy-embl/AA1981_DAT:*
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2342
1 MKYNILLILIISLFINELRA.....STTFAKFLSISLLFISFYLL 442
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                           908470 segs, 133250620 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SUMMARIES
                                                                                           OM protein – protein search, using sw model
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Maximum Match 100%
Listing first 45 summaries
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AAY97176
AAB25889
AAB25860
AAY97177
AAB25862
AAB25863
AAB25863
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Gapop 10.0 , Gapext 0.5
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Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine;
White spot disease; freshwater fish; immune response; infection control.
AAB25859 standard; Protein; 442 AA
                                                                                                                                                              48kD i-antigen protein sequence.
                                                                                                                                                                                                                                                                                              Ichthyophthirius multifiliis.
                                                                                                        18-DEC-2000 (first entry)
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ALIGNMENTS

99US-0118634. 99US-0122372. 99US-0124905. 04-FEB-2000; 2000WO-US02962. WO200046373-A1. 04-FEB-1999; 02-MAR-1999; 17-MAR-1999; 27-APR-1999; 10-AUG-2000.

FOUND INC.) UNIV GEORGIA RES FOUND) CORNELL RES FOUND INC. CLARK T G.) DICKERSON H W. (UYGE-) (DICK/) CLAR/) CORR

Clark TG, Dickerson HW, Lin T;

(first entry)

04-DEC-2000

AAY97176;

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This invention relates to novel i-antigen polypeptide sequences.

I antigens or immobilisation antigens ar common to a variety of
I antigens or immobilisation antigens ar common to a variety of
I wentonmental stimuli. This invention relates to i-antigens in
environmental stimuli. This invention relates to i-antigens in
chthyophthirus multifilis, a protozoan which is an obligate parasite
I chthyophthirus multifilis, a protozoan which is an obligate parasite
of freshwater fish causing ichthyophthiriasis or white spot disease. The
controlling induced the proposition are sequences for two
antibodies capable of binding to the nucleotide sequences and a method
antibodies capable of binding to the nucleotide sequences.
Controlling I multifilis serotypes using the nucleotide sequences.
Controlling I multifilis infection in fish. Polynucleotide or protein
controlling I multifilis infection in fish. Polynucleotide or protein
controlling I multifilis infection in fish. Sequences AAA97042,
preventing I multifilis infection in fish. Sequences AAA97042,
controlling I multifilis infection in fish. Sequences AAA97042.
Controlling I multifilis infection in fish. Sequences AAA97043.
Controlling AAAA97060) and AAA97081 represent rainigen genes and gene
certification of the invention. Sequences AAA97043. A97064
certuding AAAA97060) and AAA97071-A97088 represent primers used in the
certure controlling AAA97060 and AAA97071-A97088 represent primers used in the
certure of the invention. Sequences AAA970895999 and
contact of the invention of the sequences AAB99909 and
contact of the invention of the sequences AAB99089 and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNKNDSPAT 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EANGNOPFAANNAARGICVPCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVTDVF 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 isolation of the i-antigen gene sequences. Sequences AAB25859-B251
AAB25893-B25906 represent i-antigen protein and peptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 2342; DB 21;
100.0%; Pred. No. 2.2e-171;
Live 0; Mismatches 0;
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AAY97176
ID AAY97176 standard; Protein; 442 AA.
                                                                                                                                                                                                     Claim 1; Figure 1; 144pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 442 AA;
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Tetrahymena thermophila expresses two major beta-tubulin genes (BTU1 and ETG BTU2), which encode identical beta-tubulin proteins. Bither of these two genes (but not both at once) can be disrupted without a detectable change genes (but not both at once) can be disrupted without a detectable change in the cell phenotype. A K350L substitution in the BTU1 beta-tubulin confers increased resistance to microtubule-depolymerizing drug. CC fragment, eliminating the parallele can be transformed to pacilitaxel cells carrying the BtU1-1K350M wile a wild-type BTU1 gene fragment, eliminating the need to incorporate a means for positive containing the btU1-1K350M allele, BTU1: thermophila mutant selection. Where the bx10-1ganism is not a T. thermophila mutant selection. Where the off BTU1, can be used to generate BTU1 gene knockouts parmomycin) for that of BTU1. can be used to generate BTU1 gene knockouts parmomycin) for that of BTU1, can be used to generate BTU1 gene knockouts confaining the BLU1-1K350M allele, BTU1; and be inserted into a BTU gene by encoding antigenic polypeptides) can be inserted into a BTU gene by encoding antigenic polypeptides) can be inserted into a BTU gene by encoding antigenic polypeptides) can be inserted into a BTU gene by cuccessful cell-surface expression that is maintained by way of negative confaint of the proposus recombination-mediated insertion of a heterologous nucleic criansgenic construction in series of protein and immune response in a vertebrate. The transgenic crianlating an immune response in a vertebrate. The transgenic containing an immune response in a vertebrate. The transgenic containing an immune response in a vertebrate. The transgenic containing an immune response in a vertebrate. The transgenic containing an immune response in a vertebrate are settled on their surface multifilis immobilization-antigen (1-antigen) protein on their surface are effection by I. multifilis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Recombinant expression systems for expressing heterologous nucleic acids and producing recombinant protein, comprises nonpathogenic protozoa such as Tetrahymena resistant to paclitaxel
                                                                                                                        BTU1; beta-tubulin; protein expression system; negative selection; paclitaxel sensitivity; cell surface; antigen; protozoa; ciliate; live vaccine; Ichthyophthilus multifillis; immobilization-antigen; i-antigen; freshwater; fish; protozoacide.
                                                                                                                                                                                                                                                                                                               /note= "Gln encoded by CAR, TAG or TAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Clark TG;
                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 3A; 83pp; English.
                                                                                             48 kDa immobilization-antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                         Ichthyophthiius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                                                                                04-FEB-2000; 2000WO-US02966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; AAA52134, AAA52135.
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                                                                                                                                                                                                                                                                                                      Misc-difference 1..442
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DICKERSON H W.
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                                                                                                                                                                                                                                                                                                                                                                 WO200046381-A1
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17-MAR-1999;
27-APR-1999;
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(DICK/)
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Sequence

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55kD i-antigen protein of parasite isolate G5.
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white spot disease; freshwater fish; immune response; infection control.
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                                                                                                                            AAGAAAAGVAAVTSQCVPCQINKNDSPATAGAQANLATQCSTQCPTGTAIQDGVTLVFSN 300
                                                                                                                                   Gaps
                                                                                                                                                            9
                                         9
                        1 MKYNILLILIISLFINELRAVPCPDGTQTQAGLTDVGAADLGTCVNCRPNFYYNGGAAQG
                                DRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNKNDSPAT
                                                                                AGAQANLATQCSNQCPTGTVLDDGVTLVFNTSATLCVKCRPNFYYNGGSPQGEAPGVQVF
                                                                                                          SSTQCSQCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNTATQATQCLTTCPAGTVLD
                                                                                                                                                                              DGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVYAEATQKVQ
    Length 442;
               Indels
   DB 21;
                ö
100.0%; Score 2342; DB 2:
100.0%; Pred. No. 2.2e-17:
ive 0; Mismatches 0.
                                                                                                                                                                                                                                                                                         IAG48 (G1) surface protein amino acid sequence.
                                                                                                                                                                                                                                                  AAB25889 standard; Protein; 409 AA.
                                                                                                                                                                                                                                                                                                                                                                                                            FOUND INC
                                                                                                                                                                                                       CASTTFAKFLSISLLFISFYLL 442
                                                                                                                                                                                                               Lin T;
                                                                                                                                                                                                                                                                                                                         Ichthyophthirius multifiliis
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99US-0124905.
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                                                                                                                                                                                                                                                                                                                                                                                               99US-0131121
             Conservative
                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                 CORNELL RES FOUND
CLARK T G.
DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                          UNIV GEORGIA RES
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      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-506071/45.
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      Best Local Sim:
Matches 442;
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17-MAR-1999;
27-APR-1999;
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                 (CORR )
(CLAR/)
(DICK/)
                                                                                                                                                                                                                                                                                                                                                                                                           (UYGE-)
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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of hymenostomatid cilitates and their expression varies in response to environmental cilitates and their expression varies in response to controlled cilitates and their expression varies in response to controlled causing inchivyophthirias in obligate parasite of freshwater fish causing ichthyophthirias or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two invention are capable of binding to the nucleotide sequences and a method for identifying I. multifilis sortypes using the nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting a immune response in fish is useful for prophylaxis, treatment or for controlling I. multifilis infection in fish. Polynucleotide or protein antigence polypeptide obtained is also useful for treating or preventing I. multifilis infection in fish. Sequences AAA97064, and AAA97065 and AAA97089 represent I antigen gene sequences AAA97064, (excluding AAA97060) and AAA97089 represent primers used in the capable of the i-antigen gene sequences Sequences AAB3889 and AAB2889 and AAB2889 and AAB2889 and protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
Novel 1-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       80 PCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVTDVFDRSAAQCVKCKPNFYYNGG 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       140 SPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNKNDSPATAGAQANLATQCSNQCPTGT 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            200 VLDDGVTLVFNTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPC 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 AVPCPDGTQTQAGLTDVGAADLGTCVNCRPNFYYNGGAAQGEANGNQPFAANNAARGICV 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        260 QINKNDSPATAGAQANLATQCSTQCPTGTALQDGVTLVFSNSSTQCSQCIANYFFNGNFE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGKSQCLKCPVSKTTPAHAPGNTATQATQCLTTCPAGTVLDDGTSTNFVASATECTKCSA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 409;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 93.4%; Score 2188; DB 21; Best Local Similarity 100.0%; Pred. No. 1.3e-159; Matches 409; Conservative 0; Mismatches 0;
                                                                                                                    Disclosure; Figure 8; 144pp; English.
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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
the polypeptide ciliates and their expression varies in response to
controlmental stimuli This invention relates to i-antigens in
controlmental stimuli This invention relates to i-antigens in
controlled reasing ichthyophthitasis or white spot disease. The
controlled reasing ichthyophthitasis or white spot disease. The
controlled reasing ichthyophthitasis or white spot disease.

To invention includes two polypeptide and polynucleotide sequences for two
invention includes two polypeptide and polynucleotide sequences for two
contidentifying I. multifilis to the nucleotide sequences and a method
an immune response in fish is useful for prophytaxis, treatment or for
an immune response in fish is useful for prophytaxis, treatment or for
controlling I. multifilis infection in fish. Polynucleotide or protein
controlling I. multifilis infection in fish. Sequences AAA37043-A97042,
preventing I. multifilis infection in fish. Sequences and gene
controlling in multifilis infection in fish. Sequences and gene
fragments identified in the invention. Sequences AAA37043-A97042,
creating AAA97060) and AAA970708 represent i-antigen gene sequences.
creating and papersent i-antigen protein and peptide sequences.
AAB25893-B25906 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19;
   Immobilisation antigen; i-antigen; ichthyopthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         57 AAFVPGASTCTPCPQKKDAGAQPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECV 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  117 NCRINFYNENAPNENAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVT 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel i antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 MKYNILLILIISLFINELRAVPCPDGTQTQ-AGLTDVGAADLGT---CVNCRPNFYINGG 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               139; Indels 114; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ------NAARGICVPCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        39.3%; Score 921; DB 21; Length 468; 41.8%; Pred. No. 1.9e-62; Live 45; Mismatches 139; Indels 11.
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                                                                                                                                                                                                                                                                                                           (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; Figure 3; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                      Lin T;
                                                                                                                                                                                                                                                                                                                             CORNELL RES FOUND INC
                                                                                                                                                                                                                99US-0118634.
99US-0122372.
99US-0124905.
                                                                  Ichthyophthirius multifiliis
                                                                                                                                                                             04-FEB-2000; 2000WO-US02962.
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infection in fish
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Best Local Simil
Matches 214; C
                                                                                                         WO200046373-A1
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                                                                                                                                              10-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                          TG,
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293 GVTLVFSNSSTQCSQCIANYFFNG-NFEAGKSQCLKCPVSKTTPAHA-PGNTATQATQCL 350
                                                                                                                                                             351 TICPAGTVLDDGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAK 410
                                                                                                                                                                                                          234 APGVQVFRAGAAAAGVAAVTSQCVPCQINKN-DSPATAGAQANLATQCSTQCPTGTAIQD 292
                                                                                                                    273 APN---FNPG-----NSTCLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIAS 320
DVFDRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--N 175
              177 TDYVRSFTECVKCRLNFYYNGNN--GNTP----FNPG------KSOCTPCPAIKPAN 221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Recombinant expression systems for expressing heterologous nucleic acids and producing recombinant protein, comprises nonpathogenic protozoa such as Tetrahymena resistant to paclitaxel
                                                                                                                                                                                                                                                                                                                                                                                                                              BTUI; beta-tubulin; protein expression system; negative selection; paclitaxel sensitivity; cell surface; antigen; protozoa; ciliate; live vaccine; Ichthyophthius multifillis; immobilization-antigen; i-antigen; freshwater; fish; protozoacide.
                                                              222 VAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTE---CTNCAPNFYNN-----N
                                                176 DSPATAGAQANLATQCSNQCPTGTVLDDGVT--LVFNTSATLCVKCRPNFYYNGGSPQGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Gln encoded by CAR or TAA"
                                                                                                                                                                                                                                               VYAEATQKVQCASTTFAKFLSISLLFISFYLL 442
                                                                                                                                                                                                                                                              Clark TG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualiflers
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                                                                                                                                                                                                                                                                                                                                     AAY97177 standard; Protein; 468 AA
                                                                                                                                                                                                                                                                                                                                                                                                           55 kDa immobilization antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0122372.
99US-0124905.
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                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-514962/46.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB; AAA52136
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17-MAR-1999;
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Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.

chthyophthirius multifiliis.

Synthetic.

WO200046373-A1.

10-AUG-2000

Synthetic 55kD 1-antigen protein L6P.

18-DEC-2000 (first entry)

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Tetrahymena thermophila expresses two major beta-tubulin genes (BTU1 and BTU2), which encode identical beta-tubulin proteins. Either of these two genes (but not both at once) can be disrupted without a detectable change in the call phenotype. A #350L substitution in the BTU1 beta-tubulin can detectable change protein confers increased resistance to microtubule-depolymerizing drugs can discuss assistivity to paclitaxel, a microtubule-depolymerizing drugs calls carrying the Btu1-1K350M allele can be transformed to paclitaxel fragment, eliminating the need to incorporate a means for positive containing the Btu1-1K350M allele, BTU1: neol construct, which can be the coding region of the neol gene (conferring resistance to parmomycin) for that of BTU1, can be used to generate BTU1 gene knockouts containing at the Coding region of the neol gene (conferring resistance to parmomycin) for that of BTU1, can be used to generate BTU1 gene knockouts concessful cell-surface expression be inserted into a BTU1 gene by successful cell-surface expression that is maintained by way of negative concessful cell-surface expression that is maintained by way of negative condity restoring resistance to paclitaxel in the resulting acid, thereby restoring resistance to paclitaxel in the resulting conference of transgenic collated insertion of a heterologous nucleic transgenic host. Transgenic cliated protozoa are useful as live vaccines for stimulating an immune response in a vertebrate. The transgenic claimed). In particular, Tetrahymena expressing Ichthyopthhius contained in the resulting are also useful for producing polyclonal antibodies multifilis immobilization-antigen (1-antigen) protein on their surface are effective vehicles for vaccination of freshwater fish against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ------NAARGICVPCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVT 117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              118 DVFDRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--N 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DSPATAGAQANLATQCSNQCPTGTVLDDGVT--LVFNTSATLCVKCRPNFYYNGGSPQGE 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         234 APGVQVFAAGAAAAGVAAVTSQCVPCQINKN-DSPATAGAQANLATQCSTQCPTGTAIQD 292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       39.3%; Score 921; DB 21; Length 468;
41.8%; Pred. No. 1.9e-62;
Live 45; Mismatches 139; Indels 114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               411 VYAEATOKVQCASTTFAKFLSISLLFISFYLL 442
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Matches 214; Conservative
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(UYGE-) UNIV GEORGIA RES FOUND INC. (CORR) CORNELL RES FOUND INC. (CLAR/) CLARK T G.

DICKERSON H W.

(CLAR/) (DICK/) (LINT/)

99US-0122372. 99US-0124905. 99US-0131121.

99US-0118634

04-FEB-1999; 02-MAR-1999; 27-APR-1999; .7-MAR-1999;

04-FEB-2000; 2000WO-US02962.

Lin

Dickerson HW,

Clark TG,

WPI; 2000-506071/45.

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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
Chymorostomatid cilitates and their expression varies in response to
Chymorothalian multifillis, a protozoan which is an obligate parasite
Cofficeshwater fish causing ichthyophthiriasis or white spot disease. The
Invention includes two polypeptide and polymorlectide sequences for two
coffices capable of binding to the nucleotide sequences and a method
Coffice antibodies capable of binding to the nucleotide sequences and a method
Coffice antibodies capable of binding to the nucleotide sequences.
Coffice antibodies capable of binding to the nucleotide sequences.
Coffice antibodies capable of binding to the nucleotide or paper or or controlling in multifilis serotypes using the nucleotide sequence.
Coffice antigen Comparising a portion of the amplified product encoding an vaccines comprising a portion of the amplified product encoding an antigen polypeptide obtained is also useful for treating or preventing I multifiliis infection in fish. Sequences AAA97016-A97042.
Coffice antigen comprising and AAA97089 represent l-antigen genes and gene
Coffice and AAA97060, AAA97065 and AAA97081 represent primers used in the isolation of the invention. Sequences AAA97041-A97064
Coffice and AAA97060 and AAA97071-A97088 represent primers used in the anappring or an antigen gene sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel 1-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels 114; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 MKYNILLILIISLFINELRAVPCPDGTQTQ-AGLTDVGAADLGT---CVNCRPNFYYNGG 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the i-antigen gene sequences. Sequences AAB25859-B256
306 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 468;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   39.0%; Score 914; DB 21;
41.6%; Pred. No. 6.4e-62;
Live 45; Mismatches 140;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Figure 14; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 213; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  468 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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AAB25882 standard; Protein; 468

RESULT 6 AAB25882 ID AAB

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WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                          AAB25863;
                                                                                                                                                                           Sequence
                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                      61
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                                                                                                                                                                                                                                                                                                                           Immobilisation antigen; i-antigen; ichthyophthirlasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                         351 TTCPAGTVLDDGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAK 410
                                                                                                                                                                                      292
                                                                                                                          13 APN---FNPG------NSTCLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIAS 320
                                                                                                                                               350
                                                                                     233
         AAFVPGASTCTPCPQKKDAGAQPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECV 116
                                DVFDRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--N 175
                                                                 DSPATAGAQANLATQCSNQCPTGTVLDDGVT--LVFNTSATLCVKCRPNFYYNGGSPQGE
                                                                                                                 234 APGVQVFAAGAAAAGVAAVTSQCVPCQINKN-DSPATAGAQANLATQCSTQCPTGTALQD
                                                                                                                                               293 GVTLVFSNSSTQCSQCIANYFFNG-NFEAGKSQCLKCPVSKTTPAHA-PGNTATQATQCL
                                                                                                  222 VAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTE---CTNCAPNFYNN-----N
                         -- NAARGICVPCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVT
                                                                                                                                                                                                                                                                                                              repeat amino acid sequence SEQ ID 9.
                                                                                                                                                                                                              VYAEATQKVQCASTTFAKFLSISLLFISFYLL 442
                                                                                                                                                                                                                       Disclosure; Figure 5a; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       UNIV GEORGIA RES FOUND INC.
CORNELL RES FOUND INC.
CLARK T G.
DICKERSON H W.
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                                                                                                                                                                                                                                                                   AAB25862 standard; Protein; 89
                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                      Ichthyophthirius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                                   04-FEB-2000; 2000WO-US02962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HW,
                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dickerson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                  48kD i-antigen
                                                                                                                                                                                                                                                                                                                                                                     WO200046373-A1
                                                                                                                                                                                                                                                                                                                                                                                                                   04-FEB-1999;
02-MAR-1999;
17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                                                                                                                   18-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clark TG,
                                                                                                                                                                                                                                                                                    AAB25862;
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(CLAR/)
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                                                                                                                                                                                                                 411
              57
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pase; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                    106 CPTGTALDDGVTDVFDRSAAQCVKCKPNFYINGGSPQGEAPGVQVFAAGAAAAGVAAVTS 165
                                                                                                                                                                                                                                                                                                                                                            9
                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                               ;
                                                                                                                                                                                                                                                                                         Length 89;
                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           48kD i-antigen repeat amino acid sequence SEQ ID 10.
                                                                                                                                                                                                                                                                                         20.3%; Score 475; DB 21;
100.0%; Pred. No. 3.7e-29;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.0%; Pred. No. 3.7
Matches 89; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                        166 OCVPCQLNKNDSPATAGAQANLATQCSNQ 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYGE-) UNIV GEORGIA RES FOUND INC.
(CORR) CONNELL RES FOUND INC.
(CLAR/) CLARK T G.
(DICK/) DICKERSON H W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB25863 standard; Protein; 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ichthyophthirius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0118634.
99US-0122372.
99US-0124905.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-FEB-2000; 2000WO-US02962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0131121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     white spot disease;
                                                                                                                                                                                                                                                                       89 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         [mmobilisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200046373-A1.
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02-MAR-1999;
17-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         clark TG,
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us-09-497-967-6.rag

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This invention relates to novel i-antigen polypeptide sequences.

I fantigens or immobilisation antigens ar common to a variety of hymenostomatid ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in circulation and their expectation to the sability of freshwater fish causing ichthyophthiriasis or white spot disease. The of freshwater fish causing ichthyophthiriasis or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two artibodies capable of binding to the nucleotide sequences and a method of conting in multifiliis sercitypes using the nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting controlling I multifiliis infection in fish. Polynucleotide or protein antigenic i-antigen polypeptide obtained is also useful for treating or preventing I multifiliis infection in fish. Sequences AaA97064, fragments identified in the invention. Sequences AaA97064.

Cacluding AaA97060) and AAA97001-A97088 represent primers used in the calation of the 1-antigen protein and peptide sequences.

ABB25893-B25906 represent i antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine;
white spot disease; freshwater fish; immune response; infection control.
                 Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 195 CPTGTVLDDGVTLVFNTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTS 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20.2%; Score 472; DB 21; Length 89; 100.0%; Pred. No. 6.3e-29; 1ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48kD i-antigen repeat amino acid sequence SEQ ID 8.
                                                                                         Disclosure; Figure 5a; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        255 QCVPCQINKNDSPATAGAQANLATQCSTQ 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 QCVPCQINKNDSPATAGAQANLATQCSTQ 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYGE-) UNIV GEORGIA RES FOUND INC.
(CORR ) CORNELL RES FOUND INC.
(CLAR/) CLARK T G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB25861 standard; Protein; 83 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ichthyophthirius multifiliis
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Vert Local Similarity
Watches 89; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          89 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200046373-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-MAR-1999;
17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence
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Gaps

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Indels

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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of
hymenostomatid cilitates and their expression varies in response to
control cilitates and their expression varies in response to
control stimuli. This invention relates to i-antigens in
cof freshwater fish causing ichthyophthiriasis or white spot disease. The
cof freshwater fish causing ichthyophthiriasis or white spot disease. The
cof freshwater fish causing ichthyophthiriasis or white spot disease. The
cof invention includes two polypeptide and polyworlectide sequences for two
antibodies capable of binding to the nucleotide sequences and method
for identifying I. multifilis serotypes using the nucleotide sequences.

A composition (containing the i-antigen nucleotide) capable of eliciting
controlling I. multifilis infection in fish. Polymodieotide or protein
controlling I. multifilis infection in fish. Sequences AAA97042,
and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene
C fragments identified in the invention. Sequences AAA9704;
confronting in the invention. Sequences AAA9704;
conficient of the invention of containing seroid in the invention of containing seroid in the invention.

C (excluding AAA97065) and AAA9708B represent i-antigen genes and gene
containing AAA97065 and AAA9708B represent primers used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     isolation of the 1-antigen gene sequences. Sequences AAB28659-825889 and AAB2893-8259006 represent 1-antigen protein and peptide sequences.
                                                                                                                              Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immobilisation antigen; 1-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23 CPDGTQTQAGLTDVGAADLGTCVNCRPNFYYNGGAAQGEANGNQPFAANNAARGICVPCQ 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Саря
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19.3%; Score 451; DB 21; Length 83; 100.0%; Pred. No. 2.3e-27; .ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7;
0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    48kD 1-antigen repeat amino acid sequence SEQ ID 12.
                                                                                                                                                                                                      Disclosure; Figure 5a; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB25865 standard; Protein; 72 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           83 INRVGSVTNAGDLATLATQCSTQ 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 INRVGSVTNAGDLATLATQCSTQ 83
                                                             Lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19.38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ichthyophthirius multifillis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-FEB-2000; 2000WO-US02962.
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                                                         Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 83; Conservative
 (DICK/) DICKERSON H W. (LINT/) LIN T.
                                                                                          WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          83 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200046373-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-FEB-1999;
02-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-AUG-2000.
                                                       Clark TG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB25865;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 10
AAB25865
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10-AUG-2000.
                                                                                                                                                                 (CORR )
(CLAR/)
                                                                                                                                                                                                   DICK/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABB09437
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                                                                                                                                                                                                                                                                                                                       This invention relates to novel i-antigen polypeptide sequences.

1-antigens or immobilisation antigens ar common to a variety of
1-antigens or immobilisation antigens ar common to a variety of
2. I-antigens or immobilisation antigens ar common to a variety of
2. Invention inclates and their expression varies in response to
2. Invention includes two polypeptide and polynuclectide sequences for two
2. Invention includes two polypeptide and polynuclectide sequences for two
2. Invention includes two polypeptide and polynuclectide sequences and a method
3. Invention includes in multifilis serotypes using the nuclectide sequences.
3. Invention incontaining the insection nuclectide capable of eliciting
3. A composition (containing the insection in fish. Polynuclectide or protein
3. Controlling I. multifilis infection in fish. Polynuclectide or protein
3. Controlling I. multifilis infection in fish. Sequences and gene
3. Controlling I. multifilis infection in fish. Sequences and gene
3. Controlling I. multifilis infection in fish. Sequences and gene
3. Controlling I. multifilis infection in fish. Sequences and gene
3. Controlling I. multifilish infection in fish. Sequences and gene
4. Conduction of the invention of the antigen genes and gene
5. Controlling I. multifilish infection in fish. Sequences Analysof on the conduction of the antigen genes and gene
5. Conduction of the invention of Sequences Analysof on the conduction of the con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  fragments identified in the invention. Sequences AAA97043-A97064 (excluding AAA97066) and AAA97071-A97088 represent primers used in the (solation of the i-antigen gene sequences. Sequences AAB228859-B25889 and AAA875893-B25906 represent i antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                     Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                353 CPAGTVLDDGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVY 412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16.2%; Score 379; DB 21; Length 72; 100.0%; Pred. No. 6.5e-22; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       48kD i-antigen repeat amino acid sequence SEQ ID 11.
                                                                                                                                                                                                                                                                                                Disclosure; Figure 5a; 144pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB25864 standard; Protein; 69 AA.
                                                            (UYGE-) UNIV GEORGIA RES FOUND INC. (CORR ) CORNELL RES FOUND INC.
                                                                                                                                                              Lin T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ichthyophthirius multifiliis
            99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 100.
nes 72; Conservative
                                                                                                                                                                Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           413 AEATOKVOCAST 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 AEATQKVQCAST 72
                                                                                                              DICKERSON H W.
                                                                                                                                                                                                WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200046373-A1.
                                                                                              CLARK T
              17-MAR-1999;
                              27-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB25864;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                 TG,
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                                                                                              (CLAR/) (DICK/) I
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                                                                                                                                                                 Clark
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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
the properties and their expression varies in response to
the environmental stimuli This invention relates to i-antigens in
environmental stimuli This invention relates to i-antigens in
the protocoan which is an obligate parasite
ichthyophthrius multifilis, a protocoan which is an obligate parasite
the includes two polypeptide and polynucleotide sequences for two
invention includes two polypeptide and polynucleotide sequences for two
invention of 48 and 55 kb. Also included in the invention are
i-antigens, of 48 and 55 kb. Also included in the invention are
i-antigens, of binding to the nucleotide sequences of
antibodies capable of binding to the nucleotide sequences
for invention in multifilis is useful for prophylaxis, treatment or for
an immune response in fish is useful for prophylaxis, treatment or for
controlling in multifilis infection in fish. Polynucleotide or protein
vaccines comprising a portion of the amplified product encoding an
vaccines comprising a portion of the amplified product encoding an
vaccines comprising a polypeptide obtained is also useful for treating or
antigenic i-antigen polypeptide obtained is also useful for treating or
fragments identified in the invention. Sequences AAA37036-A97042,
certuining AAA97060, and AAA97089 represent primers used in the
certuing AAA97060, and AAA97089 represent primers used in the
certuing AAA97060, and AAA97089 represent primers used in the
certuing AAA97060, and AAA97089 represent primers used in the
certuing AAA97060, and AAA97089 represent primers used in the
certuing AAA97060, and AAA97089 represent primers used in the
certuing AAA97060, and AAA97080 represent primers used in the
certuin and papersent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            284 CPTGTAIQDGVTLVFSNSSTQCSQCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNTA 343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16.1%; Score 376; DB 21; Length 69; 100.0%; Pred. No. 1e-21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DXR; reductoisomerase; enzyme; non-mevalonate isoprenoid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            H. influenzae DXR related polypeptide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Figure 5a; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABB09437 standard; Protein; 1588 AA.
                                                                                                                                                                                                                                              GEORGIA RES FOUND INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lin
                                                                                                                                                                                                                                                                             CORNELL RES FOUND INC
                                                                                                       99US-0122372.
99US-0124905.
99US-0131121.
04-FEB-2000; 2000WO-US02962.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUL-2002 (first entry)
                                                                         99US-0118634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                             DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 69; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  344 TOATOCLIT 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        69 AA;
                                                                                                                                                                                                                                                                                                                                                                                                            (LINT/) LIN T.
                                                                                                   02-MAR-1999;
17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                                                                             UYGE-) UNIV
                                                                            04-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABB09437;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clark TG,
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The invention relates to modulating an activity of a DXR reductoisomerase enzyme of Haemophilus influenzae, comprising contacting the enzyme with a compound that modulates nor-mevalonate isopremoid biosynthesis — with a compound that modulates nor-mevalonate isopremoid biosynthesis — with a compound of menaquinone or ubiquinone. Compounds of the invention act as infected with H. influenzae having DXR reductisomerase enzyme, e.g. a infected with H. influenzae having DXR reductisomerase enzyme, e.g. a treating ear infections, conjunctivitis, menigitis, pleural empema, endocarditis and conjunctivitis, bacteraemia, sinusitis, pleural empema, endocarditis and reductisomerase enzyme related polypeptide sequence.

Soldictis The current sequence contains within it the amino acid sequence onsetting that appears to be accidentally inserted into the sequence.

Consisting the DXR encoding DNR sequence represented as an amino acid
                                                                                                                             241...1431
//note= "region that appears to be accidentally inserted
into the sequence, consisting the DXR encoding DNA
sequence represented as an amino acid sequence in three
letter code"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modulating Haemophilus influenzae DXR reductoisomerase enzyme activity, useful for treating mammals or tissues infected with H. influenzae (e.g. ear infections or pneumonia) by contacting the enzyme with a modulator of its activity
             conjunctivitis;
menaquinone; ubiquinone; virucide; ear infection; conjunctivit:
meningitis; pneumonia; conjunctivitis; bacteraemia; sinusitis;
pleural empyema; endocarditis; epiglottitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Yan K;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Jaworski DD, Payne DJ, Slater-Radosti CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 40-44; 44pp; English.
                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                             (SMIK ) SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                           09-AUG-2000; 2000US-223909P.
                                                                                                                                                                                                                                                                                                         09-AUG-2001; 2001WO-US24950.
                                                                      Haemophilus influenzae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-241698/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1588 AA;
                                                                                                                                                                                                                                 WO200211673-A2
                                                                                                                                                                                                                                                                     14-FEB-2002
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                                                                                                            Key
Region
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The present invention describes proteins and their fragments (I) encoded 20 by chromosome 2 of the human malarial parasite, Plasmodium falciparum. Also described are: (1) nucleotide sequences (II) encoding (I); and (2) vaccines against P. falciparum infection comprising (I) or (II).

(I) and (II) are useful for the development of vaccines against P. falciparum infection. (I) and polyclonal antisera or a monoclonal antibody raised to immunogens comprising the sequences of (I), are useful in the detection of infection with P. falciparum Furthermore, (I) (especially when they are rifins or secreted or membrane proteins) can aid the identification of drugs to treat or prevent P. falciparum (C) infection, or they can be used to identify drug resistance in P. falciparum. Sequencing of the Plasmodium chromosome 2 and the subsequent identification of proteins encoded by it will help to expand our understanding of parasite biology, a process hampered by the complexity of the parasite ilequence, and provide new targets for vaccine and drug development. Parasite resistance to drugs and mosquito resistance to insecticides have led to a resurgence of malaria in many
                                                                                                         24;
                                                                                                                                                                                                                                                                                                                                                                                                       651 TATTGATGCCGTAAAAACTATGGCTCGAAGCTTTTACCAGTAGATAGTGAACATAATGC 710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           598 AAACGGGTATTACTGGCAAATAAAGAATCACTGGTAACCTGC-----GGACAGCTTTT 650
                                                                                                                                                                                                                                                                                                                        -----GNOPFAANNAARGIC------VPCQINRVGSVTNAG-DLATLATQC 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          103 STQCPTGTALDDGVTDVFDRSAAQCVKCKPNFYYNGGSPQGEAFGVQVFAAGAAAAGVAA 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         711 TAICTITCAAICAIT-----ACGGC-----AGAAGCACAAAAAAATGGGTIT 755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             163 VTSQCVPCQLNKNDSPATAG-----AQANLATQCSNQCPTGTVLDDG-----VTL 207
                                                                                                                                                     20 AVPCPDGTQT-----QAGLTDVGAADL----GT----CVNCRPNFYYNGGAAQGEAN-- 63
                                                                                            Indels 116;
9.4%; Score 219; DB 23; Length 1588; 26.4%; Pred. No. 5.3e-08; Live 14; Mismatches 219; Indels 116;
                                                                          Matches 125; Conservative
                                   Local Similarity
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Query Match

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859 CAAGCGGTTGCACACCC---TAATTGGTAAAAAAAAAAATTTCTGT--CGATTCAG 913
208 VENTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQINKNDSP 267
                                            268 ATAGAQANLATQCSTQCPTGTAIQDGVTLVFSNSSTQCSQCIANYFFNGNFEAGKSQCLK 327
                                                                                         CPVSKTT----PAHAPG---NTATQATQCLTT----CPAGTVLDDGTSINFVASATEC-- 374
                                                                                                                                                                                                                                                                                                               Plasmodium falciparum; chromosome 2; human malaria parasite; vaccine; antimalarial; malaria; protozoacide; infection; insecticide.
                      816 CCGTTACACGCC----ACTTGAA-----CAATTCACCAACATAACACA----GAA
                                                                                                               CTACAATGATGAATAAGGGCTTGGAATACATTGAGGCTCGCT---GGCTTTTCAATGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Proteins encoded by chromosome 2 of the human malarial parasite, Plasmodium falciparum, useful as antimalarial vaccines and in the diagnosis of P.falciparum infection -
                                                                                                                                                           971 GIGC-GGAAGAATGGAAGT-TATTATTCATCCACAATCA-ATTATTCATTCTA 1021
                                                                                                                                     375 -TKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVYABATQKVQCASTTFA 427
                                                                                                                                                                                                                                                                                       Plasmodium falciparum chromosome 2 related protein SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gardner M, Venter JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 29-33; 577pp; English.
                                                                                                                                                                                                                    AAB18144 standard; Protein; 1700 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                          98US-0107131.
                                                                                                                                                                                                                                                                                                                                                                                                                    99WO-US26796
                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Carucci D,
                                                                                                                                                                                                                                                                                                                                                Plasmodium falciparum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-365347/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HOFF/) HOFFMAN S.
(CARU/) CARUCCI D.
(GARD/) GARDNER M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VENTER J C.
                                                                                                                                                                                                                                                                                                                                                                      WO200025728-A2.
                                                                                                                                                                                                                                                                  07-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                   05-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                         05-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                             11-MAY-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hoffman S,
                                                                                                                                                                                                                                              AAB18144;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (VENT/)
                                                                                                                                                                                                         AAB18144
ID AAB1
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RESULT 15
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                                                                                                                                                                                             16;
  parts of the world, and there is a pressing need for vaccines and new drugs. AAA10078 to AAA70287 and AAB18144 to AAB18352 represent nucleotide and protein sequences given in the present invention, but which are not specifically mentioned within the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                              703 --AATATGATT-ATAATACTAATGTTACTCATGGTTTTGGTCAAGAGTATCCTTGTGAAA 759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        249 VAAVTSQCVPCQINKNDSPATAGAQANLATQCSTQ-----CPTGTAIQDGVTLV-FSNSS 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 867 IGCGT-----TAG-----AAATTTGGAAAATATCAATGATATAGT----A 903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           363 ISTNFVASATECTKCSAGFFASKTTGFTA----GTDTCTECTKKLTSGATAKVYAEATQ 417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  866
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            362
                                                                                                                                                                                                                                                                                                                     80 PCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVTDVFDRSAAQCVKCKPNFYYNGG 139
                                                                                                                                                                                                                                                                           CIGAATIGCAIGGIGAITIGIC------AAAAGCAACAIAICCAAAIGAIAAACA 664
                                                                                                                                                                                                                                   CPDGTQTQAGLTDVGAADLGTCVNCRPNFYINGGAAQG----EANGNQPFAANNAARGICV 79
                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                               665 TCCTGAAGGATCAACAGAAATAATCCATG-------CAAACTTC-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        191 CSNQCPT-GTV-LDDGVTLVFNTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGGACATAGTAGAACGTTTTTCTGATACAGAA-----GGAGCACAATGTGATAAGAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AATAAA-----AGATAATAGTGAAGGAGCTTGCGCTCCATATAGACGATTACATGTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TQCSQCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNTATQATQCLTTCPAGTVLDDG
                                                                                                                                                                                                97;
                                                                                                                                                    DB 21; Length 1700;
                                                                                                                                                                                                16; Mismatches 202; Indels
                                                                                                                                                                                                                                                                                                                                                                                                         140 SPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNKNDSPATAGAQANLAT-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  55kD i-antigen amino acid repeat sequence SEQ ID 60.
                                                                                                                                                        9.3%; Score 218.5; DB 226.2%; Pred. No. 6.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYGE-) UNIV GEORGIA RES FOUND INC.
(CORR ) CORNELL RES FOUND INC.
(CLAR/) CLARK T G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               72 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ichthyophthirius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2000; 2000WO-US02962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB25888 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                              Best Local Similarity 26.2
Matches 112; Conservative
                                                                                                                     1700 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           964 AATCAAT 970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      418 KVQCAST 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200046373-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-FEB-1999;
02-MAR-1999;
17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB25888;
                                                                                                                       Sequence
                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           260
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                                                                                                                                                                                                                                                23
                                                                                                                                                                                                                                                                                        919
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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
hymenostomatid ciliates and their expression varies in response to
controlmental stimuli This invention relates to i-antigens in
controlmental multifilis, a protozoan which is an obligate parasite
ichthyophthrius multifilis, a protozoan which is an obligate parasite
cof freshwater fish causing ichthyophthriniasis or white spot disease. The
controlling in multifilis serotypes using the invention are
i-antipodies capable of binding to the nucleotide sequences for two
antibodies capable of binding to the nucleotide sequences of
controlling in multifilis serotypes using the nucleotide sequences.
A composition (containing is useful for prophylaxis, treatment or for
an immune response in fish is useful for prophylaxis, treatment or for
controlling in multifilis infection in fish. Polynucleotide or protein
cantigent i-antigen polypeptide obtained is also useful for treating or
antigent i-antigen polypeptide obtained is also useful for treating or
fragments identified in the invention. Sequences AAA97043-A97042,
controlling in multifilis infection in fish. Sequences and gene
fragments identified in the invention. Sequences AAA97043-A97044
cectuding AAA97060) and AAA970789 represent i-antigen gene sequences.
Cectuding AAA97060 and AAA97078 sequences. Sequences AAB28859-B25889 and
AAB25893-B25906 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human presenilin enhancer protein pen-1B derived protein SEQ ID NO: 25.
                                                                                                                                                                                                                      Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          353 CPAGTVLDDGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVY 412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              use; rat; cow; presenilin enhancer protein; pen; pen-1; pen-1B; pen-2; Aph-2; amyloid beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9.1%; Score 212; DB 21;
13.6%; Pred. No. 4.1e-09;
ve 12; Mismatches 20;
                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA014246 standard; Protein; 925 AA.
                                                                                                  ÷
                                                                                                     Lin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-MAY-2000; 2000US-0568942.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      53.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 53.63
Matches 37; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; fruit fly; mouse;
Alzheimer's disease; pen
                                                                                                     HW,
DICKERSON H W.
                                                                                                         Dickerson
                                                                                                                                                                           WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               413 AEATQKVQC 421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | : :||
61 ESAKKNIQC 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       72 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200185912-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-MAY-2002
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                                        (LINT/) LIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                         Clark TG,
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us-09-497-967-6.rag

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19;
                                                                                                                                                                                                                 The present invention relates to a method of detecting compounds capable of altering the interaction between a presentlin enhancer protein (such as pen.1, pen.18, pen.2 and Aph.2) and presentlin. The inhibition of presentlin activity causes the production of amyloid beta to be reduced and and thus be used in the treatment of Alzheimer's disease. The present exemplification of the invention.
                                                                                                               Presenilin enhancer proteins and polynucleotides useful for modulating presenilin function and screening for an agent that modulates the interaction of the protein to a binding target .
                                                             McGrath GJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            205 GAAATATCTGCTGATCTTTGGAGCGTTTGTCTCTGTGTATC--CA-----AGAA 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96 ATLATQC---STQCPTGTALDDGVTDVFDRSAAQCVKCKPNFYXNGGSPQGEA-PGVQVF 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        152 A----AGAAAAGVAAVTSQCVPCQLNKNDSPATAGAQANLATQCSNQCPTGTVLDDGVTL 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           314 AACCCAGGTGAGAGAGCACCCTC-----TATGCGA--CTGCTGGCCTAT----GTT- 358
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 208 VENTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQINKNDSP 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 359 -----TCTGGCTTGGGCTTTGGAATCATGAGTGGAGTATTTTCC----- 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    268 ATAGAQANLATQCSTQCPTGTAIQDGVTLVFSNSSTQCS------QCIANYFFNGN 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      398 TITGIGAATACCCTAIC-IGAC-----ICCTIGGGGCGAGGCACAGIGGGCAITCAIGG- 450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        318 FEAGKSQCLKCPVSKTTPAHAPGNTATQATQCLTTCPAGTV-LDDGTSTNFVASATECTK 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAADLGTCVNCR-PNFYYNGGAAQGEANGNQPFAANNAARGICVPCQINRVGSVTNAGDL 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       451 -- AGATICICCICAAITCITCCITIAITCAGCITICAIGACGCIGGICAITAICIIGCIG 508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         377 CSAGFFAS-----KTTGFTAGTDTCTECTKKLT---SGATAKVYABATQKVQCAS 423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             87;
                                                                                                                                                                                                                                                                                                                                                                                         Query Match

8.9%; Score 208.5; DB 23; Length 925;
Best Local Similarity 25.9%; Pred. No. 1.7e-07;
Matches 108; Conservative 15; Mismatches 207; Indels 87;
                                                             Nicoll SM,
                                                         Ruddy DA,
                                                                                                                                                                                  Disclosure; Page 67-70; 78pp; English.
                                                      Ellis MC,
                                                  Curtis DT, Francis GR,
                  (EXEL-) EXELIXIS INC.
                                                                                 WPI; 2002-062245/08
                                                                                                                                                                                                                                                                                                                                                            925 AA;
                                                                                                                                                                                                                                                                                                                                                            Seguence
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Search completed: February 11, 2003, 19:46:19 Job time : 35.0571 secs

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